IN THE SPECIFICATION

Please amend the title as follows:

~~NOVEL~~ POLYPEPTIDES INVOLVED IN IMMUNE RESPONSE

Please replace the Abstract with the replacement Abstract included herewith.

Please amend the paragraphs beginning on page 10, line 31 and ending on page 11, line 21 as follows:

Figures 1~~.~~–A.1-1A.2) (SEQ ID NOS: 1&2) DNA and amino acid sequence of murine CRP1 (mCRP1). Predicted signal sequence of CRP1 is underlined at the amino-terminus and the experimentally determined pro-peptide cleavage site is indicated by an asterisk. Predicted transmembrane sequence is underlined toward the carboxy-terminus. Figure 1B) (SEQ ID NOS: 3,4, 5)~~. . .~~ Amino acid alignment of B7RP1 protein sequence (mCRP1) with murine CD28.

Figures 2~~.~~–A.1-2A.2) (SEQ ID NOS: 6 & 7) DNA and amino acid sequence of murine B7RP1 (mB7RP1). Predicted signal sequence of B7RP1 is underlined at the amino-terminus and the experimentally determined pro-peptide cleavage site is indicated by an asterisk. Predicted transmembrane sequence is underlined toward the carboxy-terminus. Figure 2B) (SEQ ID NOS: 8, 9 & 10) Amino acid alignment of B7RP1 protein sequence (mB7RP1) with murine CD80 (mCD80).

Figures 3~~.~~–A.1-3A.2) (SEQ ID NOS: 11 & 12) Structure and sequence of the protein coding region of the putative human B7RP1 (hB7RP1). Predicted signal sequence of hB7RP1 is underlined at the amino-terminus. Predicted signal peptide cleavage sites are marked by asterisks. Predicted transmembrane sequence is

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underlined toward the carboxy-terminus. Figure 3B) (SEQ ID NOS: 13, 14 &15) Amino acid alignment of the putative mature hB7RP1 protein with the mature murine B7RP1 (mB7RP1) protein.

Please amend the paragraphs beginning on page 15, line 29 and ending on page 16, line 9 as follows:

Figures 12.–A.1-12A.2) (SEQ ID NOS: 16 &17) Structure and sequence of the protein coding region of human B7RP1 (hB7RP1). Predicted signal sequence of hB7RP1 is underlined at the amino-terminus. Predicted signal peptide cleavage sites are marked by asterisks. Predicted transmembrane sequence is underlined toward the carboxy-terminus. Figure 12B) (SEQ ID NOS: 18, 19 & 20) Amino acid alignment of the putative mature hB7RP1 protein (human) with the mature murine B7RP1 (mB7RP1) protein (mouse).

Figures-13.–A.1-13A.2) (SEQ ID NOS:21 &22) Structure and sequence of the protein coding region of human CRP1 (hCRP1). Predicted signal sequence of hCRP1 is underlined at the amino-terminus. Predicted signal peptide cleavage sites are marked by asterisks. Predicted transmembrane sequence is underlined toward the carboxy-terminus. Figure 13B) (SEQ ID NOS: 23 &24) Amino acid alignment of the hCRP1 protein with the murine CRP1 (~~mB7RP1~~mCRP1) protein.

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